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CLUSTAL W 2.0 multiple sequence alignment


Q5VV13|Q5VV13_HUMAN             MALSFSLLMAVLVLSYKSICSLGCDLPQTHSLGNRRALILLGQMGRISPFSCLKDRHDFR 60
P01571|IFN17_HUMAN              MALSFSLLMAVLVLSYKSICSLGCDLPQTHSLGNRRALILLAQMGRISPFSCLKDRHDFG 60
Q14CS4|Q14CS4_HUMAN             MALSFSLLMAVLVLSYKSICSLGCDLPQTHSLGNRRALILLAQMGRISHFSCLKDRHDFG 60
P01567|IFNA7_HUMAN              MARSFSLLMVVLVLSYKSICSLGCDLPQTHSLRNRRALILLAQMGRISPFSCLKDRHEFR 60
P01568|IFN21_HUMAN              MALSFSLLMAVLVLSYKSICSLGCDLPQTHSLGNRRALILLAQMGRISPFSCLKDRHDFG 60
Q5VV12|Q5VV12_HUMAN             MALSFSLLMAVLVLSYKSICSLGCDLPQTHSLGNRRALILLAQMGRISHFSCLKDRYDFG 60
P01563|IFNA2_HUMAN              MALTFALLVALLVLSCKSSCSVGCDLPQTHSLGSRRTLMLLAQMRKISLFSCLKDRHDFG 60
Q86UP4|Q86UP4_HUMAN             ----------------------MCDLPQTHSLGSRRTLMLLAQMRRISLFSCLKDRHDFG 38
Q5VYQ3|Q5VYQ3_HUMAN             MALTFYLLVALVVLSYKSFSSLGCDLPQTHSLGNRRALILLAQMRRISPFSCLKDRHDFE 60
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Q5VV13|Q5VV13_HUMAN             IPQEEFDGNQFQKAQAISVLHEMIQQTFNLFSTEDSSAAWEQSLLEKFSTELYQQLNDLE 120
P01571|IFN17_HUMAN              LPQEEFDGNQFQKTQAISVLHEMIQQTFNLFSTEDSSAAWEQSLLEKFSTELYQQLNNLE 120
Q14CS4|Q14CS4_HUMAN             FPEEEFDGHQFQKTQAISVLHEMIQQTFNLFSTEDSSAAWEQSLLEKFSTELYQQLNDLE 120
P01567|IFNA7_HUMAN              FPEEEFDGHQFQKTQAISVLHEMIQQTFNLFSTEDSSAAWEQSLLEKFSTELYQQLNDLE 120
P01568|IFN21_HUMAN              FPQEEFDGNQFQKAQAISVLHEMIQQTFNLFSTKDSSATWEQSLLEKFSTELNQQLNDLE 120
Q5VV12|Q5VV12_HUMAN             FPQEVFDGNQFQKAQAISAFHEMIQQTFNLFSTKDSSAAWDETLLDKFYIELFQQLNDLE 120
P01563|IFNA2_HUMAN              FPQEEFGN-QFQKAETIPVLHEMIQQIFNLFSTKDSSAAWDETLLDKFYTELYQQLNDLE 119
Q86UP4|Q86UP4_HUMAN             FPQEEFGN-QFQKAETIPVLHEMIQQIFNLFSTKDSSAAWDETLLDKFYTELYQQLNDLE 97
Q5VYQ3|Q5VYQ3_HUMAN             FPQEEFDDKQFQKAQAISVLHEMIQQTFNLFSTKDSSAALDETLLDEFYIELDQQLNDLE 120
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Q5VV13|Q5VV13_HUMAN             ACVIQEVGVEETPLMNEDSILAVRKYFQRITLYLIERKYSPCAWEVVRAEIMRSLSFSTN 180
P01571|IFN17_HUMAN              ACVIQEVGMEETPLMNEDSILAVRKYFQRITLYLTEKKYSPCAWEVVRAEIMRSLSFSTN 180
Q14CS4|Q14CS4_HUMAN             ACVIQEVGVEETPLMNEDSILAVRKYFQRITLYLTEKKYSPCAWEVVRAEIMRSLSFSTN 180
P01567|IFNA7_HUMAN              ACVIQEVGVEETPLMNEDFILAVRKYFQRITLYLMEKKYSPCAWEVVRAEIMRSFSFSTN 180
P01568|IFN21_HUMAN              ACVIQEVGVEETPLMNVDSILAVKKYFQRITLYLTEKKYSPCAWEVVRAEIMRSFSLSKI 180
Q5VV12|Q5VV12_HUMAN             ACVTQEVGVEEIALMNEDSILAVRKYFQRITLYLMGKKYSPCAWEVVRAEIMRSFSFSTN 180
P01563|IFNA2_HUMAN              ACVIQGVGVTETPLMKEDSILAVRKYFQRITLYLKEKKYSPCAWEVVRAEIMRSFSLSTN 179
Q86UP4|Q86UP4_HUMAN             ACVIQGVGVTETPLMKEDSILAVRKYFQRITLYLKEKKYSPCAWEVVRAEIMRSFSLSTN 157
Q5VYQ3|Q5VYQ3_HUMAN             SCVMQEVGVIESPLMYEDSILAVRKYFQRITLYLTEKKYSSCAWEVVRAEIMRSFSLSIN 180
                                :** * **: * .**  * ****:**********  :***.*************:*:*


Q5VV13|Q5VV13_HUMAN             LQKRLRRKD 189
P01571|IFN17_HUMAN              LQKILRRKD 189
Q14CS4|Q14CS4_HUMAN             LQKRLRRKD 189
P01567|IFNA7_HUMAN              LKKGLRRKD 189
P01568|IFN21_HUMAN              FQERLRRKE 189
Q5VV12|Q5VV12_HUMAN             LQKGLRRKD 189
P01563|IFNA2_HUMAN              LQESLRSKE 188
Q86UP4|Q86UP4_HUMAN             LQESLRSKE 166
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Q5VYQ3|Q5VYQ3_HUMAN                 LQKRLKSKE 189
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